



EXHIBIT A

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Search for

☐ 1: CAA09617. serum albumin [Mu...
[gi:3647327]

BLink, Domains, Nucleotide, Related Sequences, Domain Relatives,
Taxonomy, LinkOut

LOCUS CAA09617 608 aa linear ROD 02-FEB-2000

DEFINITION serum albumin [Mus musculus].

ACCESSION CAA09617

VERSION CAA09617.1 GI:3647327

DBSOURCE embl locus MMU011413, accession [AJ011413.1](#)

KEYWORDS

SOURCE house mouse.

ORGANISM Mus musculus

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

REFERENCE 1 (residues 1 to 608)

AUTHORS Cable, J., Harris, P.D., Tinsley, R.C. and Lazarus, C.M.

TITLE Phylogenetic analysis of Gyrodactylus spp. (Platyhelminthes:
Monogenea) using rDNA sequences

JOURNAL Can. J. Zool. 77, 1439-1449 (1999)

REFERENCE 2 (residues 1 to 608)

AUTHORS Van Reeth, T., Gabant, P., Dreze, P., Szpirer, J. and Szpirer, C.

JOURNAL Unpublished

REFERENCE 3 (residues 1 to 608)

AUTHORS Van Reeth, T.

TITLE Direct Submission

JOURNAL Submitted (18-SEP-1998) Van Reeth T., Departement de Biologie
Moleculaire, Universite Libre de Bruxelles, Laboratoire de Biologie
du Developpement, rue des chevaux, 67, 1640 Rhode St Genese,
BELGIUM

FEATURES

source

Location/Qualifiers

1..608

/organism="Mus musculus"

/db_xref="taxon:10090"

/tissue_type="liver"

Protein

1..608

CDS

1..608

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/db_xref="SWISS-PROT:P07724"

ORIGIN

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121 ernecflqhk ddnpplppfe rpeaeamcts fkenpttfmg hylhevarrh pyfyapelly
181 yaeqyneilt qccaeadkes cltpkldgk ekalvssvrq rmkcsmmqkf gerafkawav
241 arlsqtfpna dfaeitklat dltkvnkecc hgdllcadd raelakymce ngatissklq
301 tcdkplllk ahclsevehd tmpadlpaia adfvedqevc knyaeakdvf lgtflyeysr
361 rhpdyvsll lrlakkyeat lekccaeapn pacygtvlae fqplveepkn lvktncdlye
421 klgeygfqna ilvrytqkap qvstptlvea arnlgrvgtk cctlpedqrl pcvedylsai
481 lnrvcllhek tpvsehvtkc csgslverrp cfsaltvdet yvpkefkaet ftfhdsictl
541 pekekqikkq talaelvkhk pkataeqlkt vmddfaqfld tckaadkdt cfstegpnlv
601 trckdala

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PubMed Nucleotide Protein Genome Structure PopSet Taxonomy OMIM Boo

Search for

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☐ 1: CAA23753. reading frame
HSA...[gi:28590]

BLink, Domains, Nucleotide, OMIM, Related Sequences, Domain Relatives,
PubMed, SNP, Taxonomy, LinkOut

LOCUS CAA23753 609 aa linear PRI 12-SEP-1993
DEFINITION reading frame HSA [Homo sapiens].
ACCESSION CAA23753
VERSION CAA23753.1 GI:28590
DBSOURCE embl locus HSALB1, accession V00494.1
KEYWORDS .
SOURCE human.
ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE 1 (residues 1 to 609)
AUTHORS Dugaiczky,A., Law,S.W. and Dennison,O.E.
TITLE Nucleotide sequence and the encoded amino acids of human serum
albumin mRNA
JOURNAL Proc. Natl. Acad. Sci. U.S.A. 79 (1), 71-75 (1982)
MEDLINE 82105994
PUBMED 6275391

FEATURES Location/Qualifiers
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/db_xref="taxon:9606"
Protein 1..609
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CDS 1..609
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/db_xref="SWISS-PROT:P02768"

ORIGIN
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121 grnecflqhk ddnplnlprlv rpevdvmcta fhdneetflk kylieiarrh pyfyapellf
181 fakrykaaft eccqaadkaa clpkldelr degkassakq rlkcaslqkf gerafkawav
241 arlsqrfpka efaevsklvt dltkvhtecc hgdlllecadd radlakyice nqdsissklk
301 eccepllek shciaevend empadlpsla adfveskdvc knyaeakdvf lgmflyeyar
361 rhpdyssvll lrlaktyett lekccaaadp hecyakvfde fkplveepqn likqncelfe
421 qlgeykfqna llvrytkkvp evstptlvev srnlgkvgs cckhpeakrm pcaedylsvv
481 lnqlcvlhek tpsvdrvtkc cteslvnrrp cfsalevdet yvpkefnaet ftfhadictl
541 sekerqikkq talvelvkhk pkatkeqlka vmddfaafve kckkaddket cfaeegkklv
601 aasqaalgl
//

Revised: October 24, 2001.

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